

## SEQUENCE LISTING

<110> Xia, Menghang  
Williams, Mark E.

<120> Nucleic Acid Molecules Encoding Novel  
Human Low-Voltage Activated Calcium Channel Proteins,  
Designated - Alpha 1I-1 and Alpha 1I-2, Encoded Proteins and  
Methods of Use Thereof

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<140> PCT/US2004/031820

<141> 2004-09-28

<150> 60/508,112

<151> 2003-10-02

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			20					25					30		
Ser	Ser	Pro	Pro	Gly	Leu	Glu	Glu	Pro	Leu	Asp	Gly	Ala	Asp	Pro	His
		35				40					45				
Val	Pro	His	Pro	Asp	Leu	Ala	Pro	Ile	Ala	Phe	Phe	Cys	Leu	Arg	Gln
	50					55				60					
Thr	Thr	Ser	Pro	Arg	Asn	Trp	Cys	Ile	Lys	Met	Val	Cys	Asn	Pro	Trp
65				70						75					80
Phe	Glu	Cys	Val	Ser	Met	Leu	Val	Ile	Leu	Leu	Asn	Cys	Val	Thr	Leu
			85					90					95		
Gly	Met	Tyr	Gln	Pro	Cys	Asp	Asp	Met	Asp	Cys	Leu	Ser	Asp	Arg	Cys
			100					105					110		
Lys	Ile	Leu	Gln	Val	Phe	Asp	Asp	Phe	Ile	Phe	Ile	Phe	Phe	Ala	Met
		115				120						125			
Glu	Met	Val	Leu	Lys	Met	Val	Ala	Leu	Gly	Ile	Phe	Gly	Lys	Lys	Cys
	130					135					140				
Tyr	Leu	Gly	Asp	Thr	Trp	Asn	Arg	Leu	Asp	Phe	Phe	Ile	Val	Met	Ala
145				150						155					160
Gly	Met	Val	Glu	Tyr	Ser	Leu	Asp	Leu	Gln	Asn	Ile	Asn	Leu	Ser	Ala
			165					170					175		
Ile	Arg	Thr	Val	Arg	Val	Leu	Arg	Pro	Leu	Lys	Ala	Ile	Asn	Arg	Val
			180					185					190		
Pro	Ser	Met	Arg	Ile	Leu	Val	Asn	Leu	Leu	Leu	Asp	Thr	Leu	Pro	Met
		195				200						205			
Leu	Gly	Asn	Val	Leu	Leu	Leu	Cys	Phe	Phe	Val	Phe	Phe	Ile	Phe	Gly
	210					215					220				
Ile	Ile	Gly	Val	Gln	Leu	Trp	Ala	Gly	Leu	Leu	Arg	Asn	Arg	Cys	Phe
225				230						235					240
Leu	Glu	Glu	Asn	Phe	Thr	Ile	Gln	Gly	Asp	Val	Ala	Leu	Pro	Pro	Tyr
			245					250					255		
Tyr	Gln	Pro	Glu	Glu	Asp	Asp	Glu	Met	Pro	Phe	Ile	Cys	Ser	Leu	Ser
			260				265						270		
Gly	Asp	Asn	Gly	Ile	Met	Gly	Cys	His	Glu	Ile	Pro	Pro	Leu	Lys	Glu
		275				280						285			
Gln	Gly	Arg	Glu	Cys	Cys	Leu	Ser	Lys	Asp	Asp	Val	Tyr	Asp	Phe	Gly
	290					295					300				
Ala	Gly	Arg	Gln	Asp	Leu	Asn	Ala	Ser	Gly	Leu	Cys	Val	Asn	Trp	Asn
305				310						315					320
Arg	Tyr	Tyr	Asn	Val	Cys	Arg	Thr	Gly	Ser	Ala	Asn	Pro	His	Lys	Gly
			325					330						335	
Ala	Ile	Asn	Phe	Asp	Asn	Ile	Gly	Tyr	Ala	Trp	Ile	Val	Ile	Phe	Gln
			340				345						350		
Val	Ile	Thr	Leu	Glu	Gly	Trp	Val	Glu	Ile	Met	Tyr	Tyr	Val	Met	Asp
		355				360						365			
Ala	His	Ser	Phe	Tyr	Asn	Phe	Ile	Tyr	Phe	Ile	Leu	Leu	Ile	Ile	Val
	370					375					380				
Gly	Ser	Phe	Phe	Met	Ile	Asn	Leu	Cys	Leu	Val	Val	Ile	Ala	Thr	Gln
385				390						395					400
Phe	Ser	Glu	Thr	Lys	Gln	Arg	Glu	His	Arg	Leu	Met	Leu	Glu	Gln	Arg
			405					410					415		
Gln	Arg	Tyr	Leu	Ser	Ser	Ser	Thr	Val	Ala	Ser	Tyr	Ala	Glu	Pro	Gly



			420					425				430				
Asp	Cys	Tyr	Glu	Glu	Ile	Phe	Gln	Tyr	Val	Cys	His	Ile	Leu	Arg	Lys	
		435					440					445				
Ala	Lys	Arg	Arg	Ala	Leu	Gly	Leu	Tyr	Gln	Ala	Leu	Gln	Ser	Arg	Arg	
	450					455					460					
Gln	Ala	Leu	Gly	Pro	Glu	Ala	Pro	Ala	Pro	Ala	Lys	Pro	Gly	Pro	His	
465					470			475							480	
Ala	Lys	Glu	Pro	Arg	His	Tyr	Gln	Leu	Cys	Pro	Gln	His	Ser	Pro	Leu	
				485				490						495		
Asp	Ala	Thr	Pro	His	Thr	Leu	Val	Gln	Pro	Ile	Pro	Ala	Thr	Leu	Ala	
		500					505					510				
Ser	Asp	Pro	Ala	Ser	Cys	Pro	Cys	Cys	Gln	His	Glu	Asp	Gly	Arg	Arg	
	515					520						525				
Pro	Ser	Gly	Leu	Gly	Ser	Thr	Asp	Ser	Gly	Gln	Glu	Gly	Ser	Gly	Ser	
	530					535					540					
Gly	Ser	Ser	Ala	Gly	Gly	Glu	Asp	Glu	Ala	Asp	Gly	Asp	Gly	Ala	Arg	
545				550				555							560	
Ser	Ser	Glu	Asp	Gly	Ala	Ser	Ser	Glu	Leu	Gly	Lys	Glu	Glu	Glu	Glu	
				565				570						575		
Glu	Glu	Gln	Ala	Asp	Gly	Ala	Val	Trp	Leu	Cys	Gly	Asp	Val	Trp	Arg	
		580					585					590				
Glu	Thr	Arg	Ala	Lys	Leu	Arg	Gly	Ile	Val	Asp	Ser	Lys	Tyr	Phe	Asn	
	595					600						605				
Arg	Gly	Ile	Met	Met	Ala	Ile	Leu	Val	Asn	Thr	Val	Ser	Met	Gly	Ile	
	610					615					620					
Glu	His	His	Glu	Gln	Pro	Glu	Glu	Leu	Thr	Asn	Ile	Leu	Glu	Ile	Cys	
625				630				635							640	
Asn	Val	Val	Phe	Thr	Ser	Met	Phe	Ala	Leu	Glu	Met	Ile	Leu	Lys	Leu	
				645				650						655		
Ala	Ala	Phe	Gly	Leu	Phe	Asp	Tyr	Leu	Arg	Asn	Pro	Tyr	Asn	Ile	Phe	
		660					665					670				
Asp	Ser	Ile	Ile	Val	Ile	Ile	Arg	Pro	Pro	Thr	Ala	Ala	Ser	Tyr	Leu	
	675					680						685				
Tyr	Pro	Gly	Pro	Ala	Leu	Arg	Asp	Arg	Ser	Ile	Trp	Glu	Ile	Val	Gly	
	690					695					700					
Gln	Ala	Asp	Gly	Gly	Leu	Ser	Val	Leu	Arg	Thr	Phe	Arg	Leu	Leu	Arg	
705				710				715							720	
Val	Leu	Lys	Leu	Val	Arg	Phe	Met	Pro	Ala	Leu	Arg	Arg	Gln	Leu	Val	
				725				730						735		
Val	Leu	Met	Lys	Thr	Met	Asp	Asn	Val	Ala	Thr	Phe	Cys	Met	Leu	Leu	
		740					745					750				
Met	Leu	Phe	Ile	Phe	Ile	Phe	Ser	Ile	Leu	Gly	Met	His	Ile	Phe	Gly	
	755					760						765				
Cys	Lys	Phe	Ser	Leu	Arg	Thr	Asp	Thr	Gly	Asp	Thr	Val	Pro	Asp	Arg	
	770					775					780					
Lys	Asn	Phe	Asp	Ser	Leu	Leu	Trp	Ala	Ile	Val	Thr	Val	Phe	Gln	Ile	
785				790				795							800	
Leu	Thr	Gln	Glu	Asp	Trp	Asn	Val	Val	Leu	Tyr	Asn	Gly	Met	Ala	Ser	
			805					810						815		
Thr	Ser	Pro	Trp	Ala	Ser	Leu	Tyr	Phe	Val	Ala	Leu	Met	Thr	Phe	Gly	
		820					825						830			
Asn	Tyr	Val	Leu	Phe	Asn	Leu	Leu	Val	Ala	Ile	Leu	Val	Glu	Gly	Phe	
	835					840						845				
Gln	Ala	Glu	Gly	Asp	Ala	Asn	Arg	Ser	Tyr	Ser	Asp	Glu	Asp	Gln	Ser	
	850					855					860					
Ser	Ser	Asn	Ile	Glu	Glu	Phe	Asp	Lys	Leu	Gln	Glu	Gly	Leu	Asp	Ser	
865				870				875							880	

Ser	Gly	Asp	Pro	Lys	Leu	Cys	Pro	Ile	Pro	Met	Thr	Pro	Asn	Gly	His	
				885					890					895		
Leu	Asp	Pro	Ser	Leu	Pro	Leu	Gly	Gly	His	Leu	Gly	Pro	Ala	Gly	Ala	
			900					905					910			
Ala	Gly	Pro	Ala	Pro	Arg	Leu	Ser	Leu	Gln	Pro	Asp	Pro	Met	Leu	Val	
		915					920					925				
Ala	Leu	Gly	Ser	Arg	Lys	Ser	Ser	Val	Met	Ser	Leu	Gly	Arg	Met	Ser	
	930					935					940					
Tyr	Asp	Gln	Arg	Ser	Leu	Ser	Ser	Ser	Arg	Ser	Ser	Tyr	Tyr	Gly	Pro	
945					950					955					960	
Trp	Gly	Arg	Ser	Ala	Ala	Trp	Ala	Ser	Arg	Arg	Ser	Ser	Trp	Asn	Ser	
				965					970					975		
Leu	Lys	His	Lys	Pro	Pro	Ser	Ala	Glu	His	Glu	Ser	Leu	Leu	Ser	Ala	
			980					985					990			
Glu	Arg	Gly	Gly	Gly	Ala	Arg	Val	Cys	Glu	Val	Ala	Ala	Asp	Glu	Gly	
		995					1000					1005				
Pro	Pro	Arg	Ala	Ala	Pro	Leu	His	Thr	Pro	His	Ala	His	His	Val	His	
	1010					1015					1020					
His	Gly	Pro	His	Leu	Ala	His	Arg	His	Arg	His	His	Arg	Arg	Thr	Leu	
1025				1030						1035					1040	
Ser	Leu	Asp	Asn	Arg	Asp	Ser	Val	Asp	Leu	Ala	Glu	Leu	Val	Pro	Ala	
			1045					1050						1055		
Val	Gly	Ala	His	Pro	Arg	Ala	Ala	Trp	Arg	Ala	Ala	Gly	Pro	Ala	Pro	
		1060						1065					1070			
Gly	His	Glu	Asp	Cys	Asn	Gly	Arg	Met	Pro	Ser	Ile	Ala	Lys	Asp	Val	
	1075					1080						1085				
Phe	Thr	Lys	Met	Gly	Asp	Arg	Gly	Asp	Arg	Gly	Glu	Asp	Glu	Glu	Glu	
	1090					1095					1100					
Ile	Asp	Tyr	Thr	Leu	Cys	Phe	Arg	Val	Arg	Lys	Met	Ile	Asp	Val	Tyr	
1105				1110						1115					1120	
Lys	Pro	Asp	Trp	Cys	Glu	Val	Arg	Glu	Asp	Trp	Ser	Val	Tyr	Leu	Phe	
			1125					1130						1135		
Ser	Pro	Glu	Asn	Arg	Phe	Arg	Val	Leu	Cys	Gln	Thr	Ile	Ile	Ala	His	
		1140						1145					1150			
Lys	Leu	Phe	Asp	Tyr	Val	Val	Leu	Ala	Phe	Ile	Phe	Leu	Asn	Cys	Ile	
	1155						1160					1165				
Thr	Ile	Ala	Leu	Glu	Arg	Pro	Gln	Ile	Glu	Ala	Gly	Ser	Thr	Glu	Arg	
	1170					1175					1180					
Ile	Phe	Leu	Thr	Val	Ser	Asn	Tyr	Ile	Phe	Thr	Ala	Ile	Phe	Val	Gly	
1185				1190						1195					1200	
Glu	Met	Thr	Leu	Lys	Val	Val	Ser	Leu	Gly	Leu	Tyr	Phe	Gly	Glu	Gln	
			1205					1210					1215			
Ala	Tyr	Leu	Arg	Ser	Ser	Trp	Asn	Val	Leu	Asp	Gly	Phe	Leu	Val	Phe	
		1220						1225					1230			
Val	Ser	Ile	Ile	Asp	Ile	Val	Val	Ser	Leu	Ala	Ser	Ala	Gly	Gly	Ala	
	1235					1240						1245				
Lys	Ile	Leu	Gly	Val	Leu	Arg	Val	Leu	Arg	Leu	Leu	Arg	Thr	Leu	Arg	
	1250					1255					1260					
Pro	Leu	Arg	Val	Ile	Ser	Arg	Ala	Pro	Gly	Leu	Lys	Leu	Val	Val	Glu	
1265				1270						1275					1280	
Thr	Leu	Ile	Ser	Ser	Leu	Lys	Pro	Ile	Gly	Asn	Ile	Val	Leu	Ile	Cys	
			1285					1290						1295		
Cys	Ala	Phe	Phe	Ile	Ile	Phe	Gly	Ile	Leu	Gly	Val	Gln	Leu	Phe	Lys	
	1300						1305					1310				
Gly	Lys	Phe	Tyr	His	Cys	Leu	Gly	Val	Asp	Thr	Arg	Asn	Ile	Thr	Asn	
	1315					1320						1325				
Arg	Ser	Asp	Cys	Met	Ala	Ala	Asn	Tyr	Arg	Trp	Val	His	His	Lys	Tyr	

1330	1335	1340
Asn Phe Asp Asn Leu Gly Gln Ala Leu Met Ser Leu Phe Val Leu Ala		
1345	1350	1355
Ser Lys Asp Gly Trp Val Asn Ile Met Tyr Asn Gly Leu Asp Ala Val		1360
	1365	1370
Ala Val Asp Gln Gln Pro Val Thr Asn His Asn Pro Trp Met Leu Leu		1375
	1380	1385
Tyr Phe Ile Ser Phe Leu Leu Ile Val Ser Phe Phe Val Leu Asn Met		1390
	1395	1400
Phe Val Gly Val Val Val Glu Asn Phe His Lys Cys Arg Gln His Gln		1405
	1410	1415
Glu Ala Glu Glu Ala Arg Arg Arg Glu Glu Lys Arg Leu Arg Arg Leu		1420
1425	1430	1435
Glu Lys Lys Arg Arg Lys Ala Gln Arg Leu Pro Tyr Tyr Ala Thr Tyr		1440
	1445	1450
Cys His Thr Arg Leu Leu Ile His Ser Met Cys Thr Ser His Tyr Leu		1455
	1460	1465
Asp Ile Phe Ile Thr Phe Ile Ile Cys Leu Asn Val Val Thr Met Ser		1470
	1475	1480
Leu Glu His Tyr Asn Gln Pro Thr Ser Leu Glu Thr Ala Leu Lys Tyr		1485
	1490	1495
Cys Asn Tyr Met Phe Thr Thr Val Phe Val Leu Glu Ala Val Leu Lys		1500
1505	1510	1515
Leu Val Ala Phe Gly Leu Arg Arg Phe Phe Lys Asp Arg Trp Asn Gln		1520
	1525	1530
Leu Asp Leu Ala Ile Val Leu Leu Ser Val Met Gly Ile Thr Leu Glu		1535
	1540	1545
Glu Ile Glu Ile Asn Ala Ala Leu Pro Ile Asn Pro Thr Ile Ile Arg		1550
	1555	1560
Ile Met Arg Val Leu Arg Ile Ala Arg Val Leu Lys Leu Leu Lys Met		1565
	1570	1575
Ala Thr Gly Met Arg Ala Leu Leu Asp Thr Val Val Gln Ala Leu Pro		1580
1585	1590	1595
Gln Val Gly Asn Leu Gly Leu Leu Phe Met Leu Leu Phe Phe Ile Tyr		1600
	1605	1610
Ala Ala Leu Gly Val Glu Leu Phe Gly Lys Leu Val Cys Asn Asp Glu		1615
	1620	1625
Asn Pro Cys Glu Gly Met Ser Arg His Ala Thr Phe Glu Asn Phe Gly		1630
	1635	1640
Met Ala Phe Leu Thr Leu Phe Gln Val Ser Thr Gly Asp Asn Trp Asn		1645
	1650	1655
Gly Ile Met Lys Asp Thr Leu Arg Asp Cys Thr His Asp Glu Arg Ser		1660
1665	1670	1675
Cys Leu Ser Ser Leu Gln Phe Val Ser Pro Leu Tyr Phe Val Ser Phe		1680
	1685	1690
Val Leu Thr Ala Gln Phe Val Leu Ile Asn Val Val Val Ala Val Leu		1695
	1700	1705
Met Lys His Leu Asp Asp Ser Asn Lys Glu Ala Gln Glu Asp Ala Glu		1710
	1715	1720
Met Asp Ala Glu Leu Glu Leu Glu Met Ala His Gly Leu Gly Pro Gly		1725
	1730	1735
Pro Arg Leu Pro Thr Gly Ser Pro Gly Ala Pro Gly Arg Gly Pro Gly		1740
1745	1750	1755
Gly Ala Gly Gly Gly Gly Asp Thr Glu Gly Gly Leu Cys Arg Arg Cys		1760
	1765	1770
Tyr Ser Pro Ala Gln Glu Asn Leu Trp Leu Asp Ser Val Ser Leu Ile		1775
	1780	1785
		1790

Ile	Lys	Asp	Ser	Leu	Glu	Gly	Glu	Leu	Thr	Ile	Ile	Asp	Asn	Leu	Ser	
		1795					1800					1805				
Gly	Ser	Ile	Phe	His	His	Tyr	Ser	Ser	Pro	Ala	Gly	Cys	Lys	Lys	Cys	
	1810					1815					1820					
His	His	Asp	Lys	Gln	Glu	Val	Gln	Leu	Ala	Glu	Thr	Glu	Ala	Phe	Ser	
1825					1830					1835					1840	
Leu	Asn	Ser	Asp	Arg	Ser	Ser	Ser	Ile	Leu	Leu	Gly	Asp	Asp	Leu	Ser	
			1845						1850					1855		
Leu	Glu	Asp	Pro	Thr	Ala	Cys	Pro	Pro	Gly	Arg	Lys	Asp	Ser	Lys	Gly	
		1860					1865						1870			
Glu	Leu	Asp	Pro	Pro	Glu	Pro	Met	Arg	Val	Gly	Asp	Leu	Gly	Glu	Cys	
	1875					1880					1885					
Phe	Phe	Pro	Leu	Ser	Ser	Thr	Ala	Val	Ser	Pro	Asp	Pro	Glu	Asn	Phe	
	1890					1895					1900					
Leu	Cys	Glu	Met	Glu	Glu	Ile	Pro	Phe	Asn	Pro	Val	Arg	Ser	Trp	Leu	
1905					1910					1915					1920	
Lys	His	Asp	Ser	Ser	Gln	Ala	Pro	Pro	Ser	Pro	Phe	Ser	Pro	Asp	Ala	
			1925						1930					1935		
Ser	Ser	Pro	Leu	Leu	Pro	Met	Pro	Ala	Glu	Phe	Phe	His	Pro	Ala	Val	
		1940						1945					1950			
Ser	Ala	Ser	Gln	Lys	Gly	Pro	Glu	Lys	Gly	Thr	Gly	Thr	Gly	Thr	Leu	
	1955					1960						1965				
Pro	Lys	Ile	Ala	Leu	Gln	Gly	Ser	Trp	Ala	Ser	Leu	Arg	Ser	Pro	Arg	
	1970				1975						1980					
Val	Asn	Cys	Thr	Leu	Leu	Arg	Gln	Val	Pro	Thr	Pro	Pro	Arg	Pro		
1985					1990					1995						